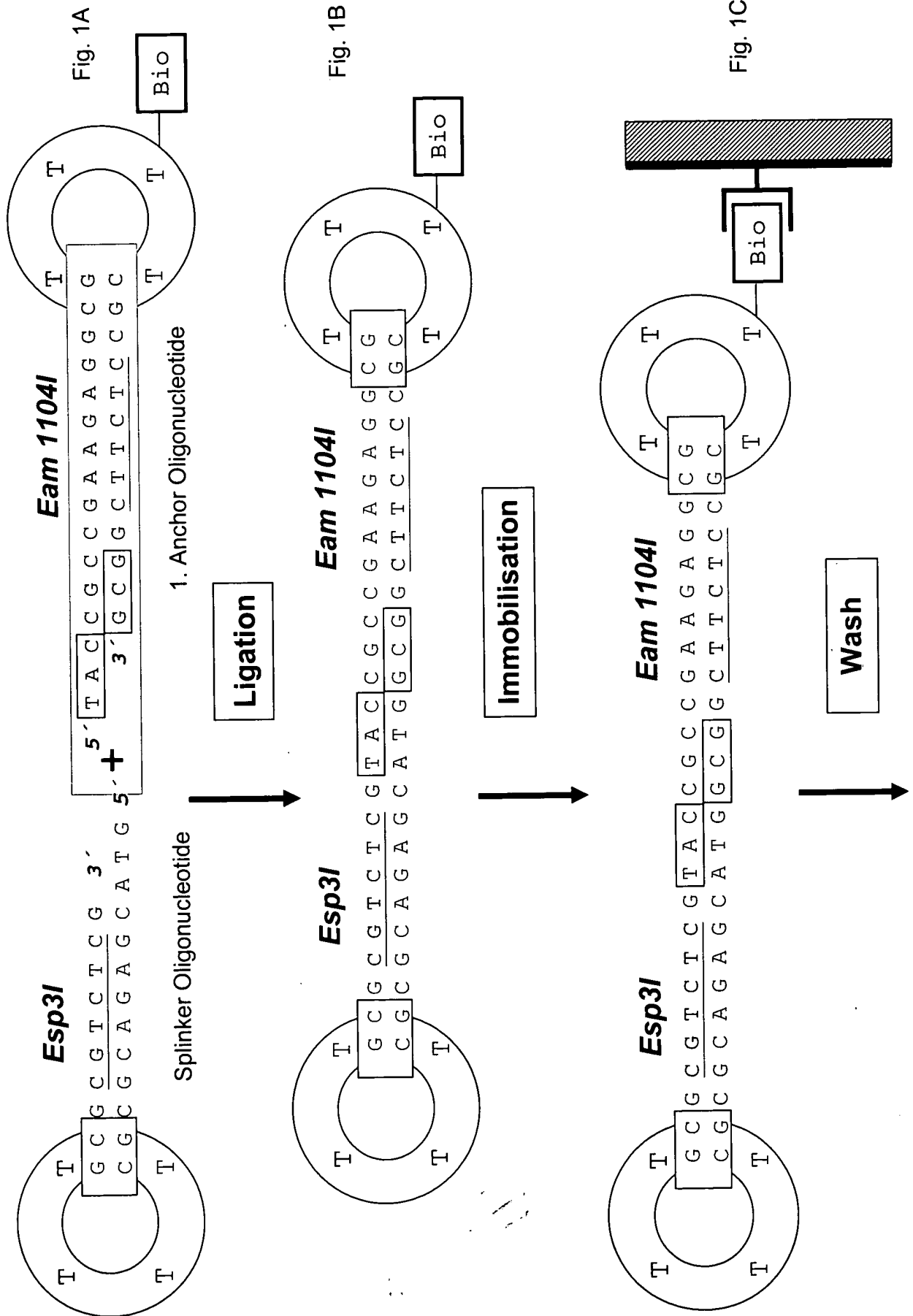
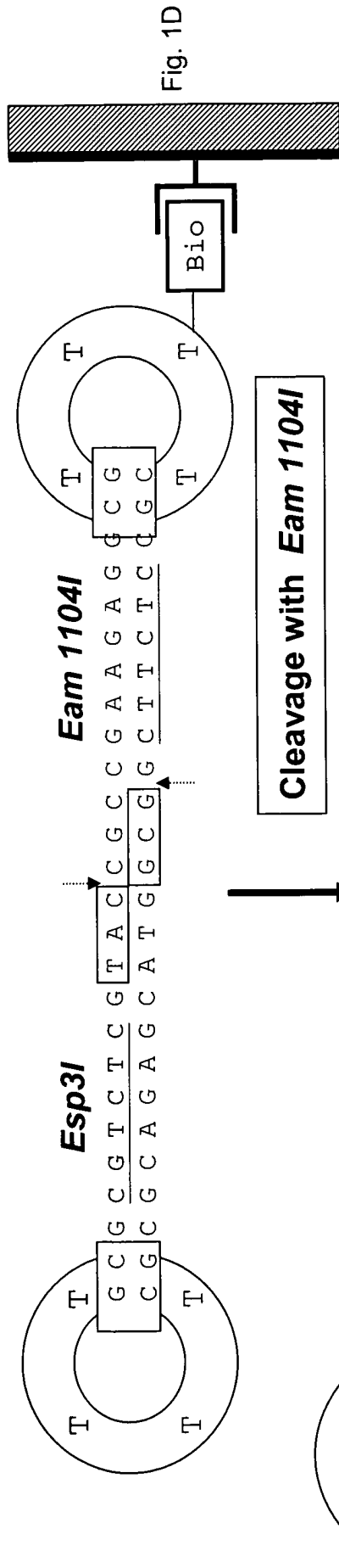


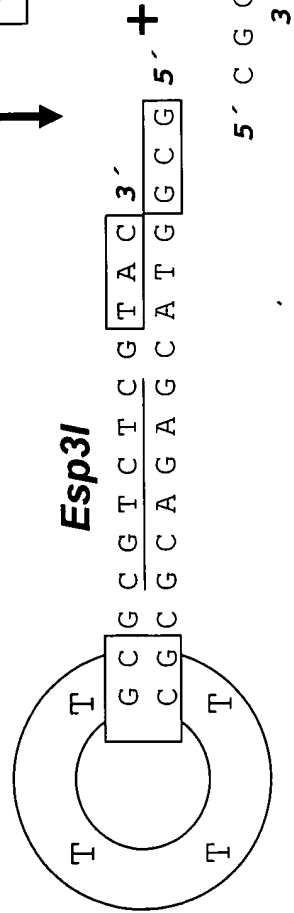
(RSPS Variant 1)

Fig. 1 3nt Overhang elongation

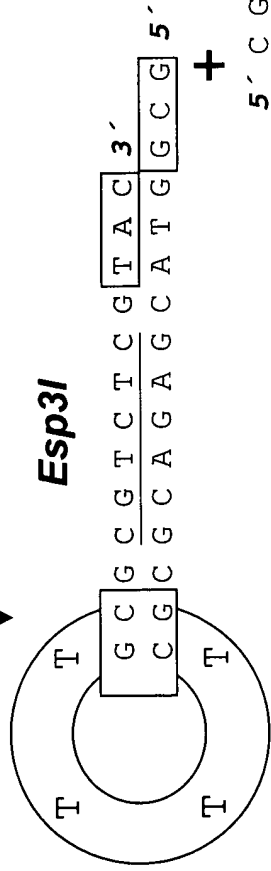




Cleavage with *Eam 1104I*



Transfer of the supernatant including the Elongated splinker to the next reaction



Elongated Splinker Oligonucleotide

Ligation

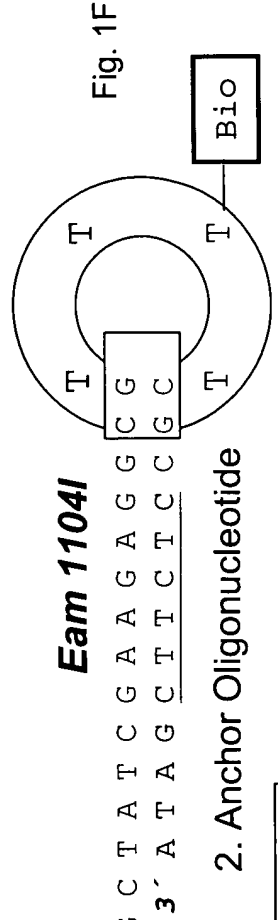
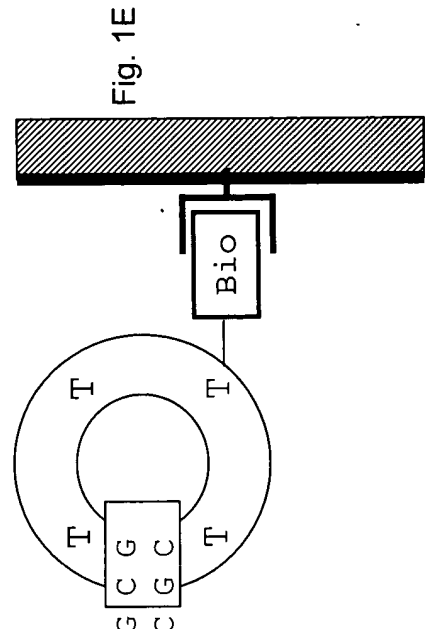


Fig. 1F



[illegible]

Fig. 3 3nt Overhang elongation

(RLPS Variant 1)

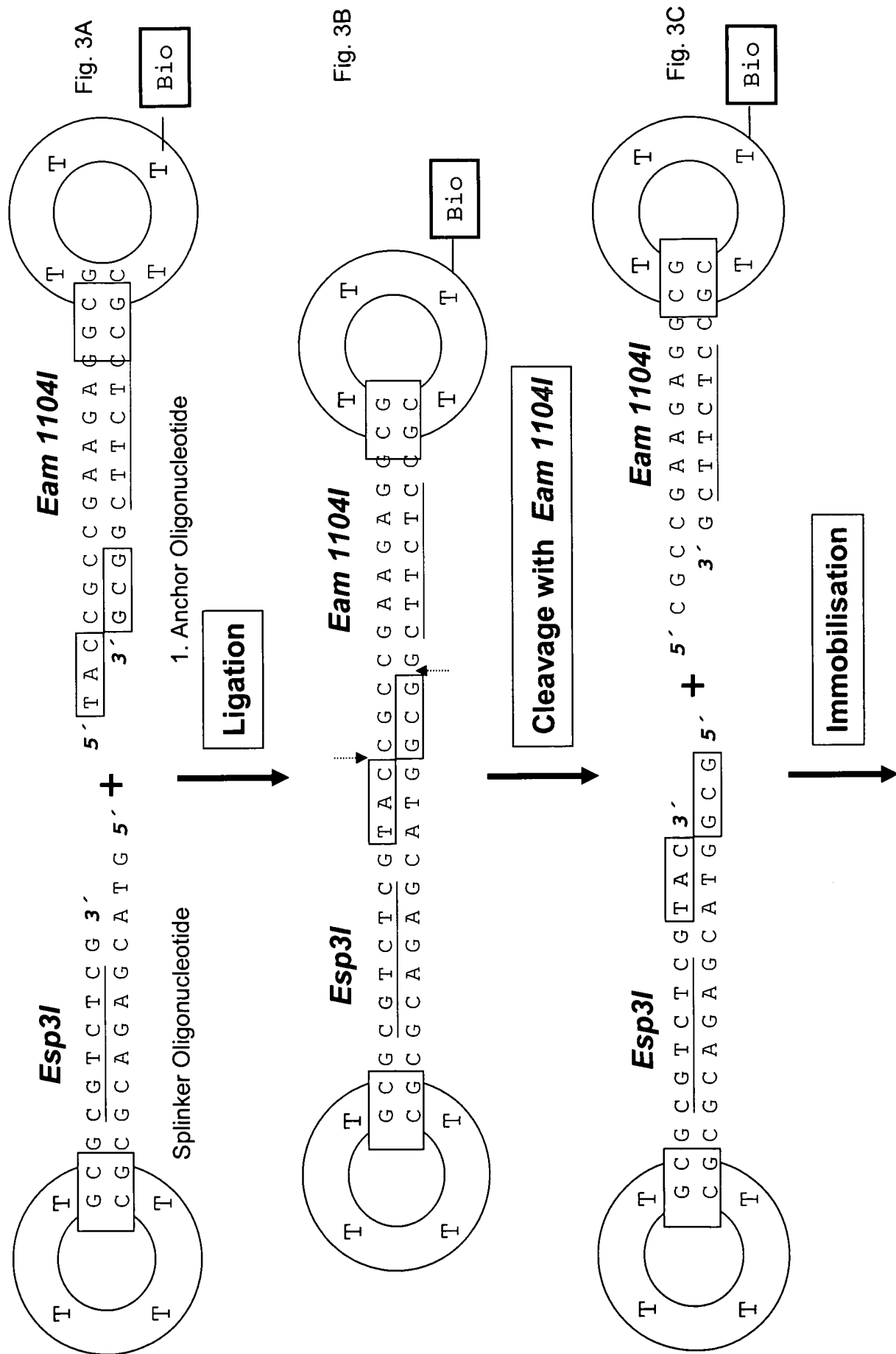


Fig. 3D

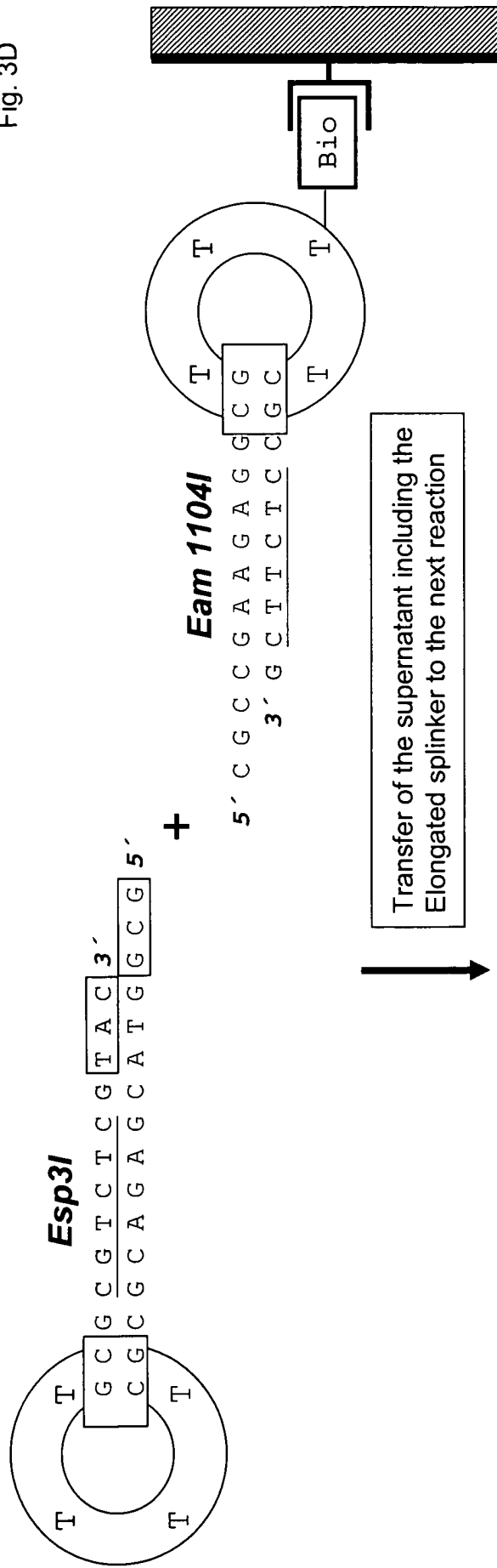


Fig. 3E

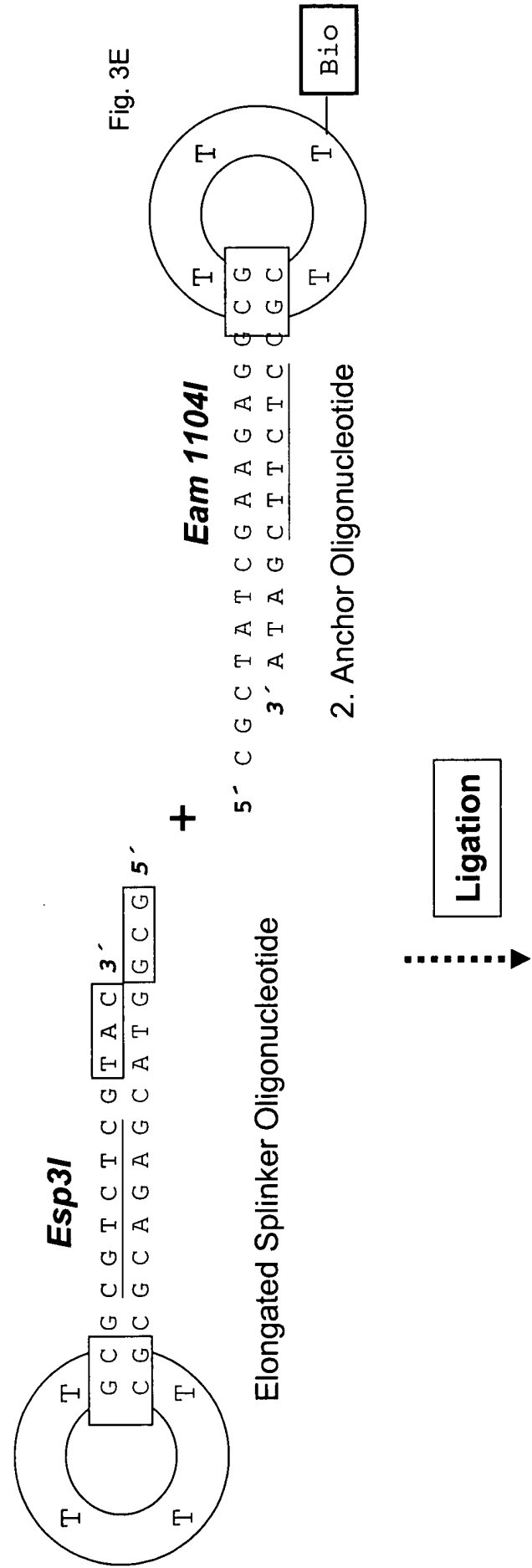


Fig. 4D

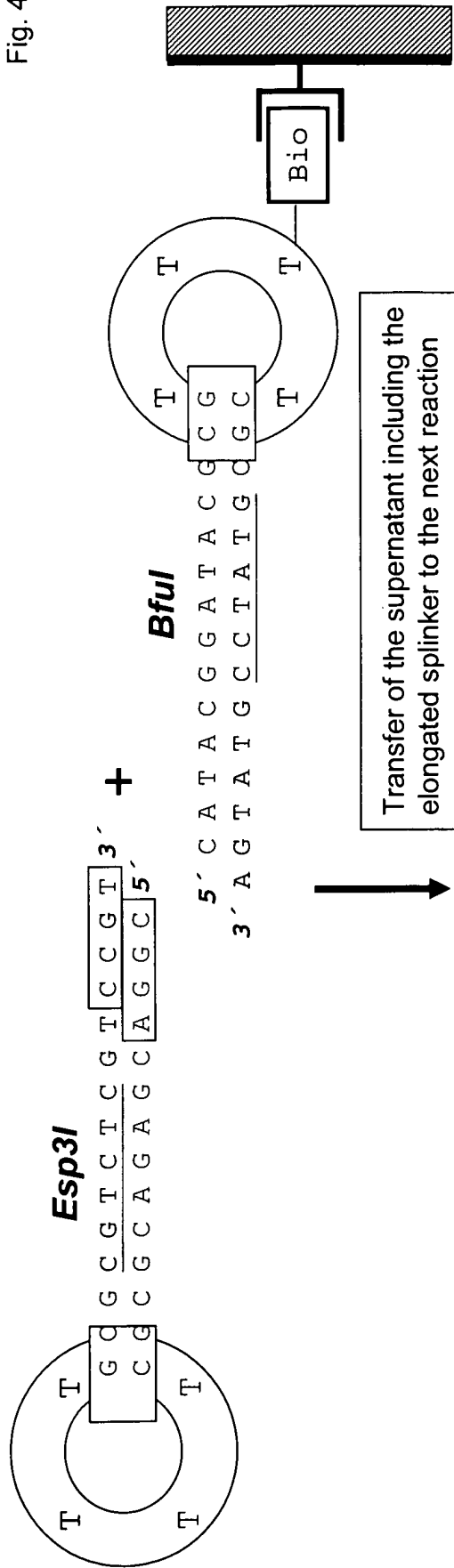


Fig. 4E

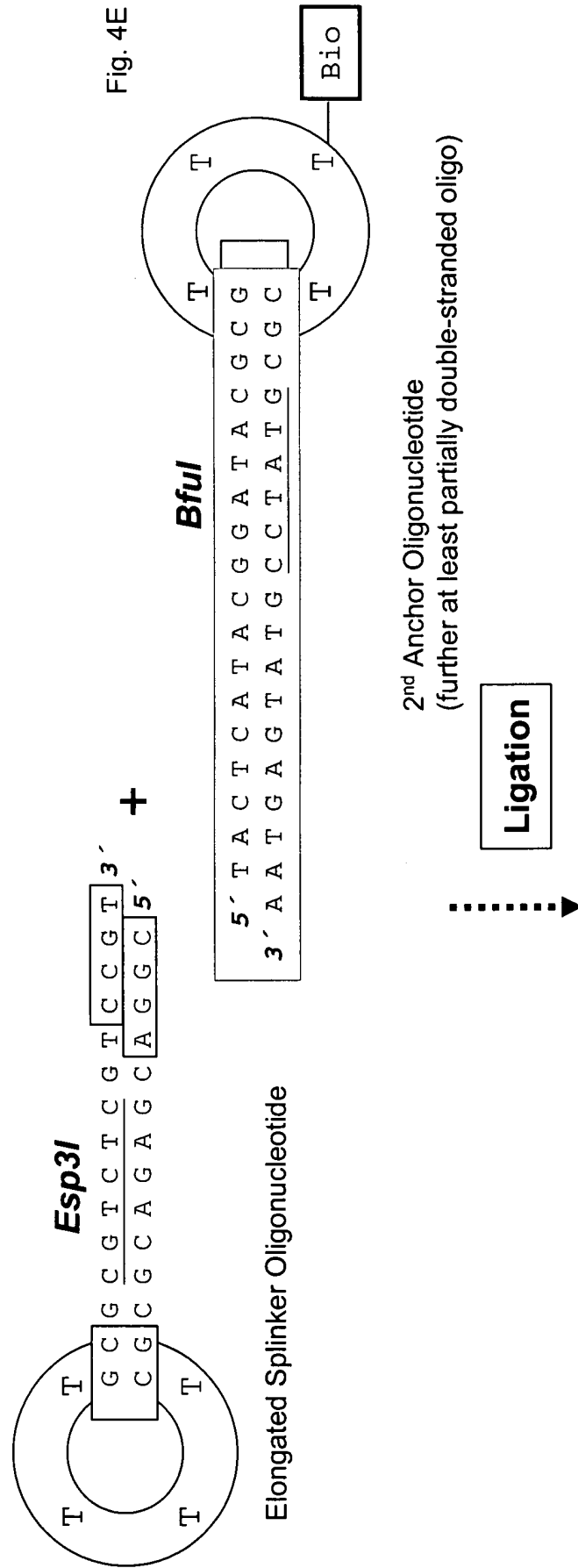
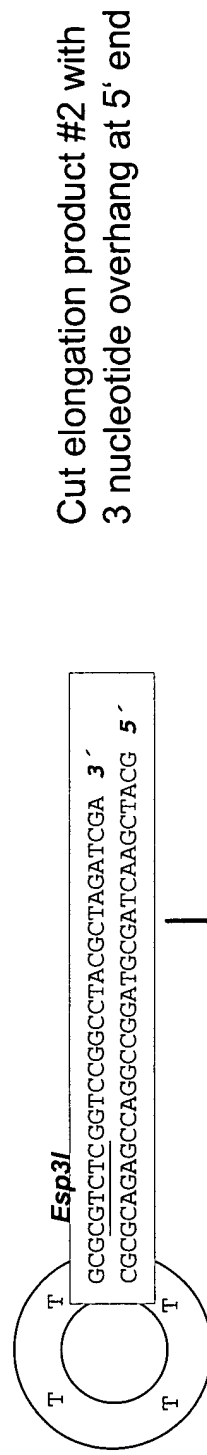
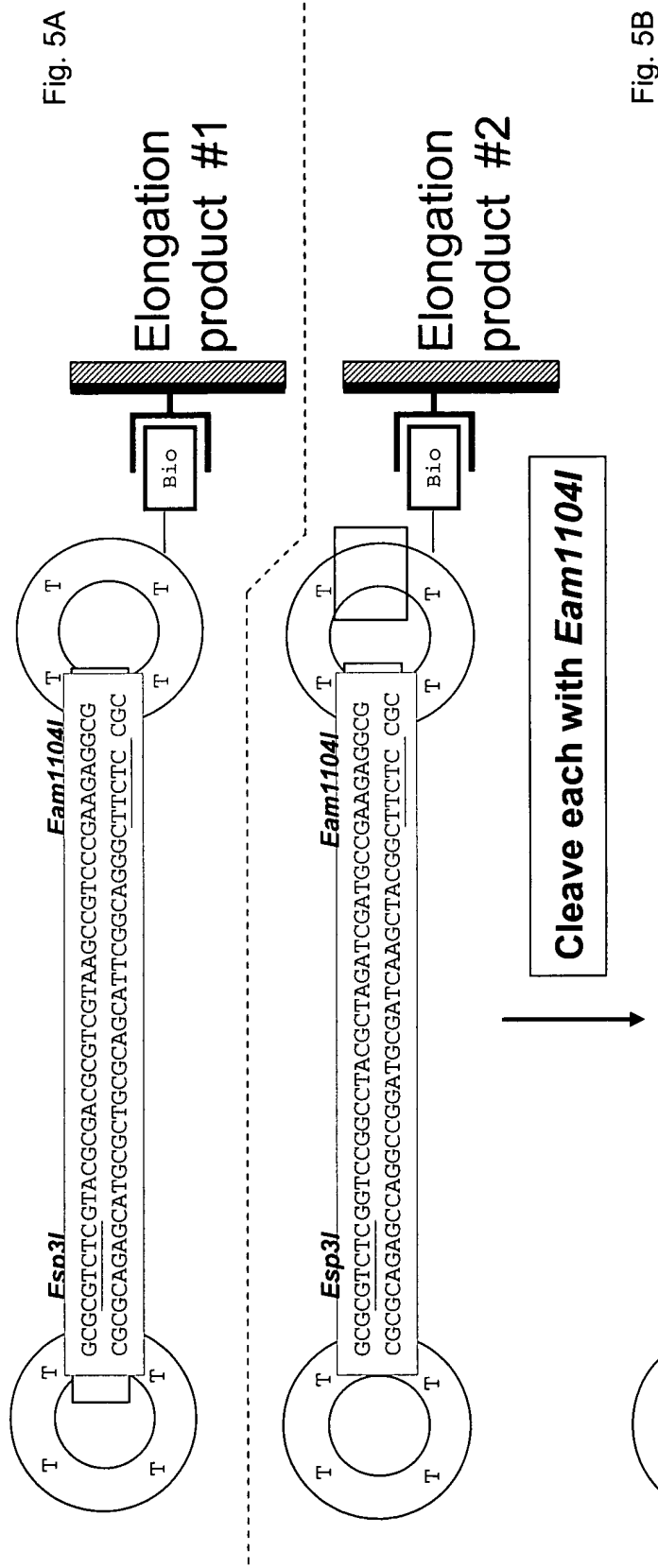
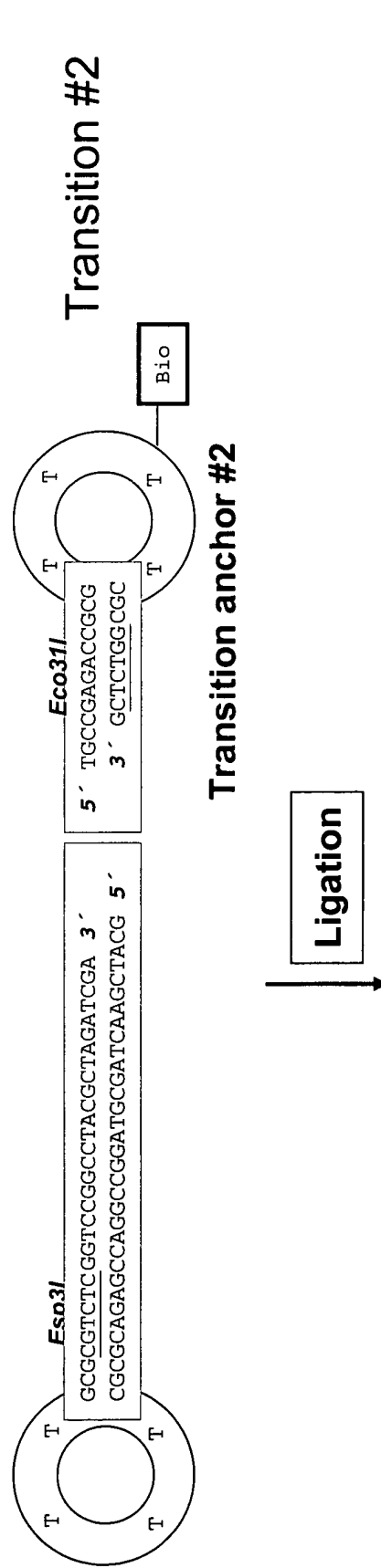
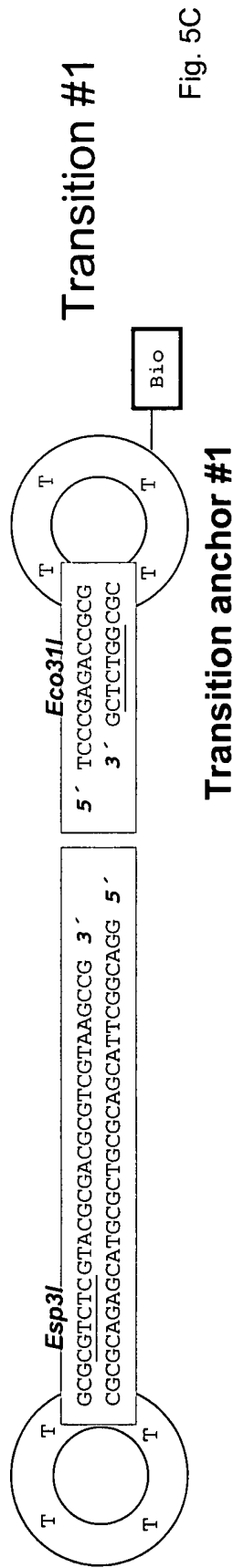
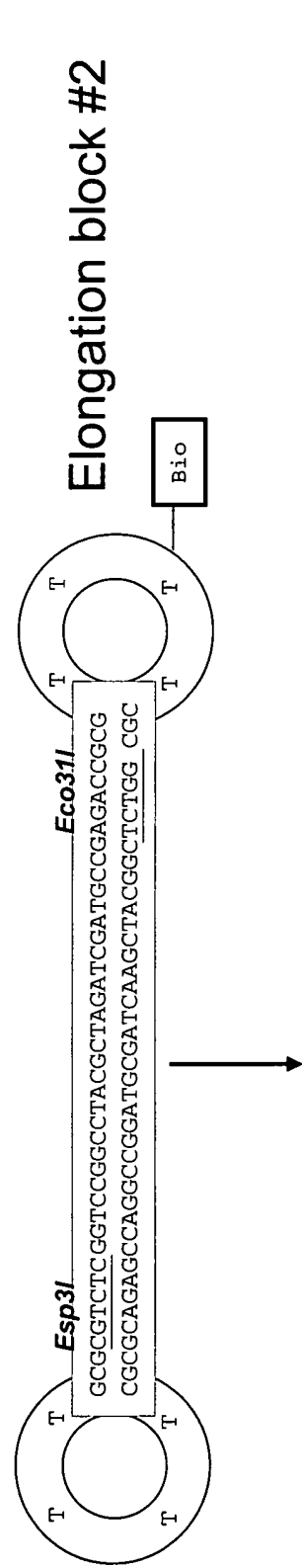
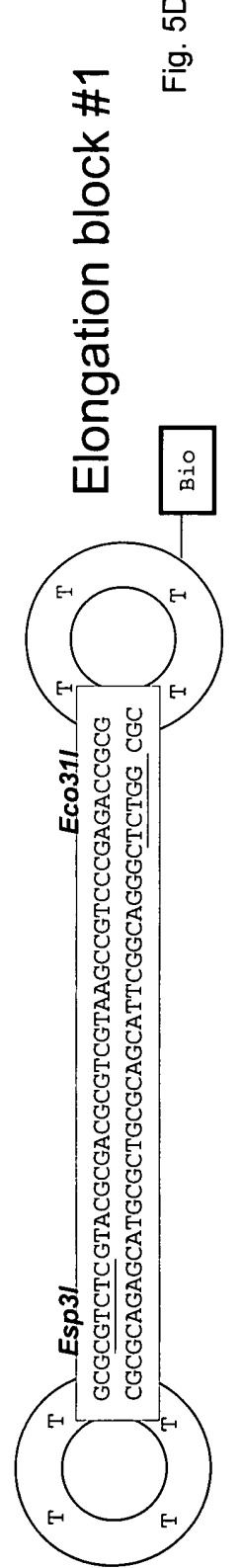


Fig. 5 Addition of transition anchor (both RSPS and RLPS) and first transposition (3 nt overhang)





Ligation



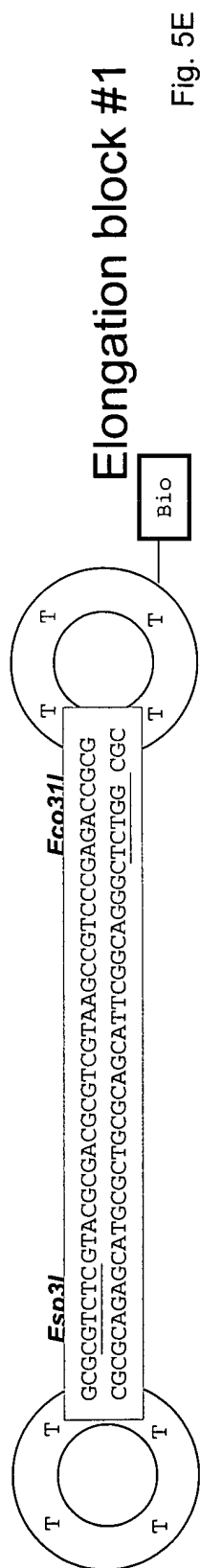
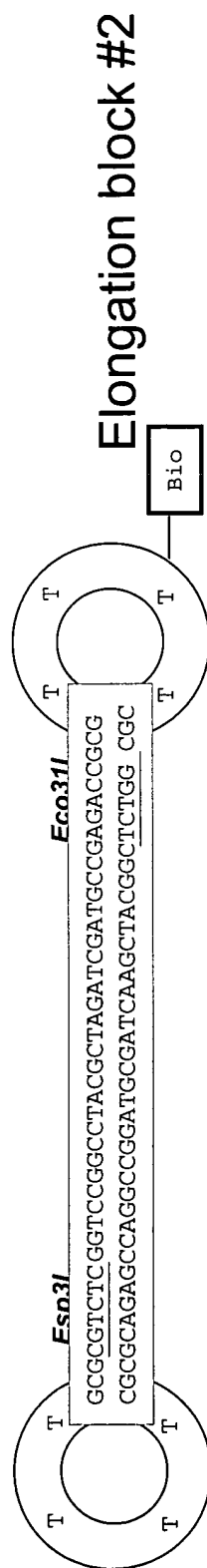
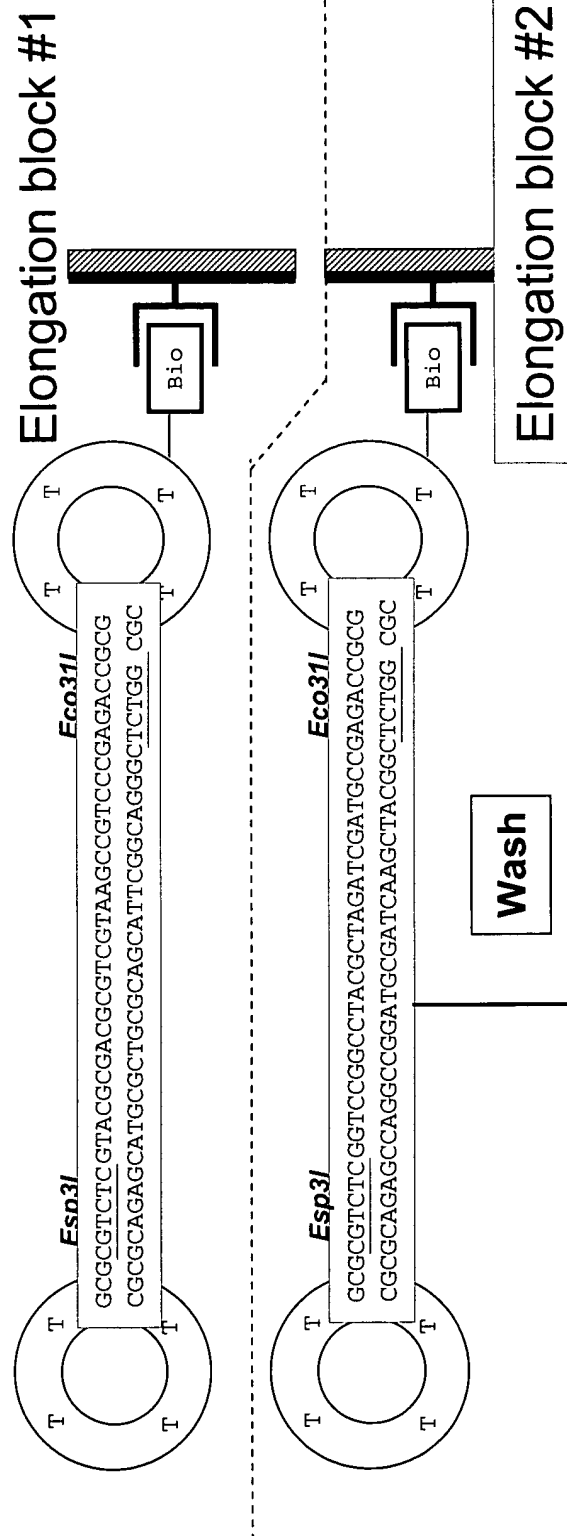


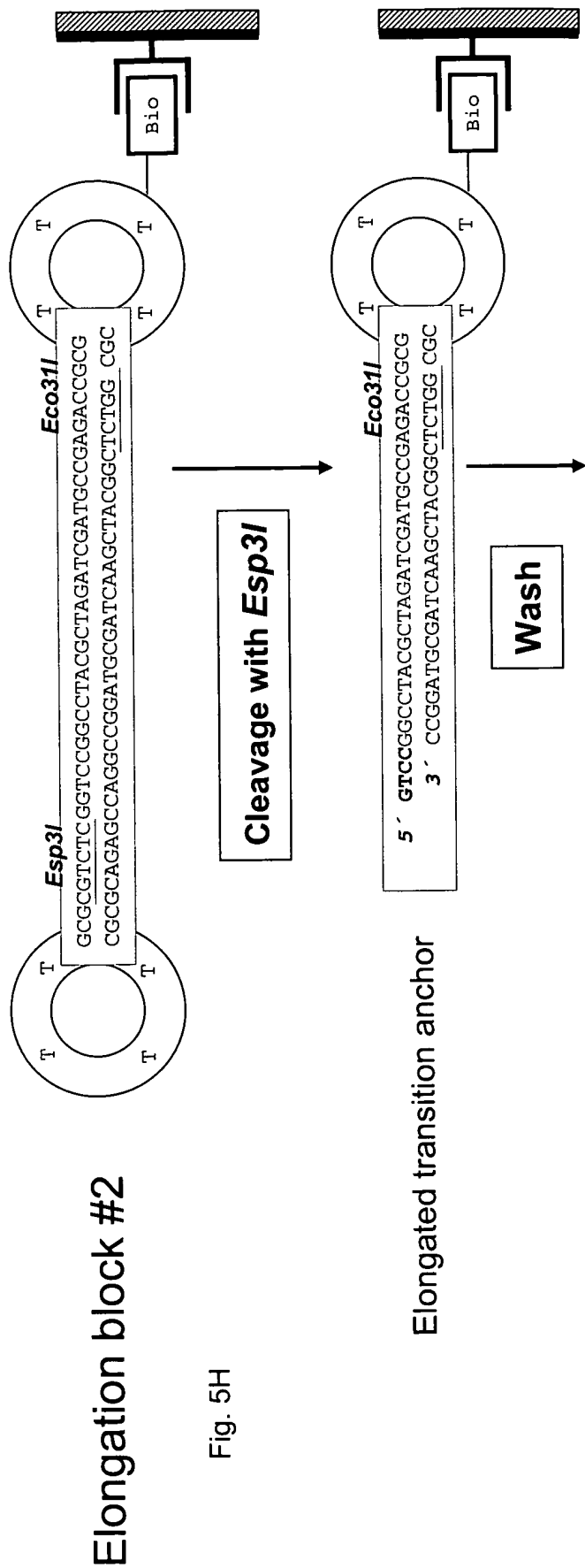
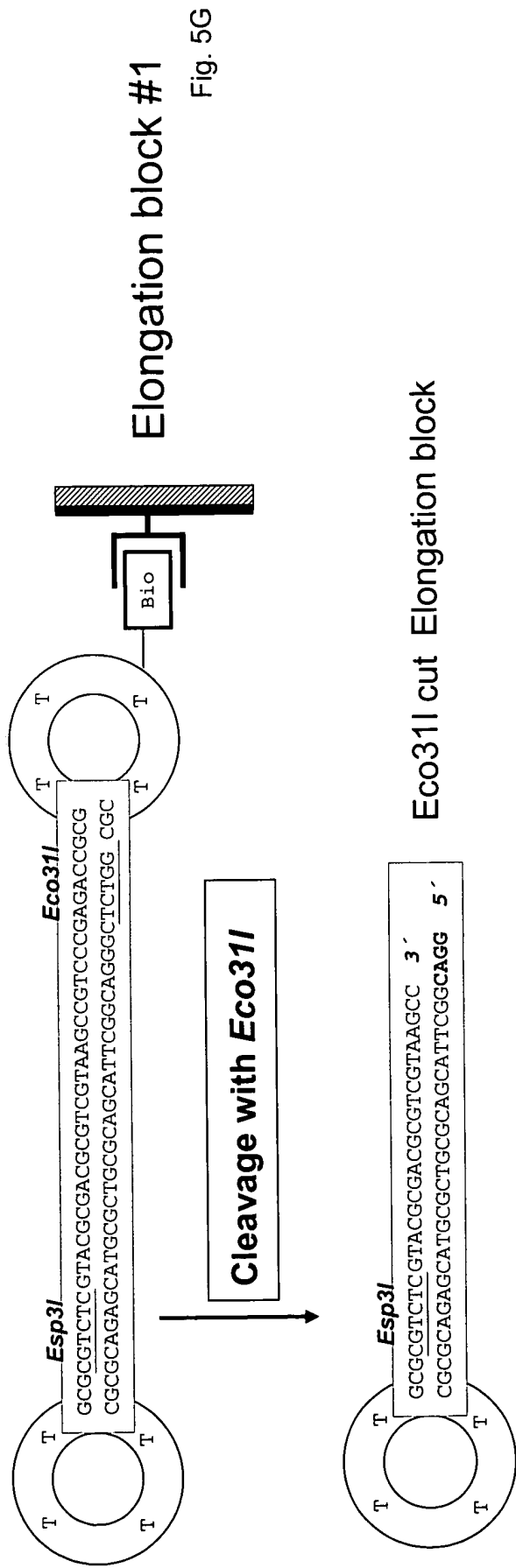
Fig. 5E



Immobilisation

Fig. 5F





Transfer supernatant with cut elongation block from elongation #1 to an elongated transition anchor

Fig. 5I

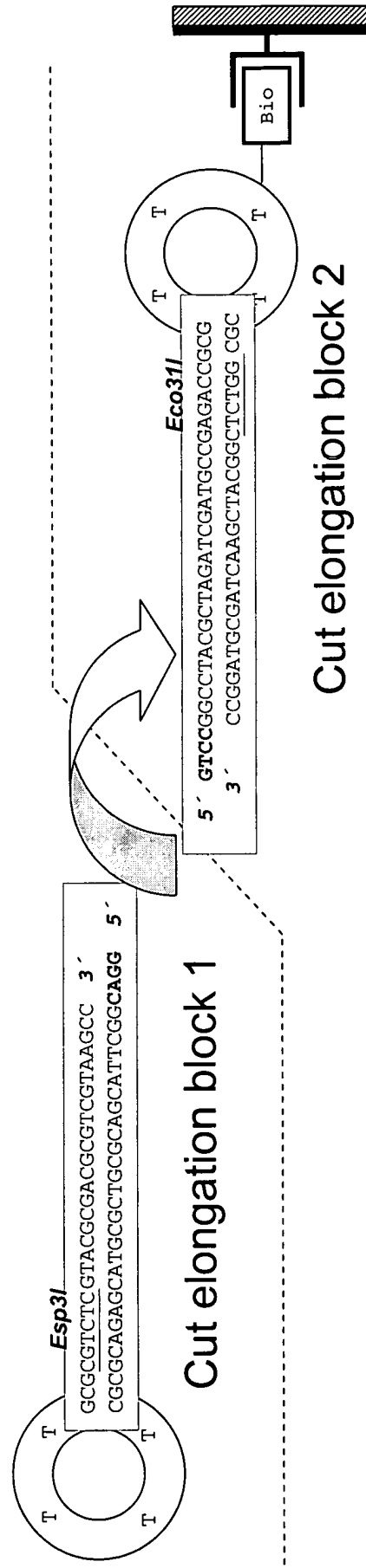
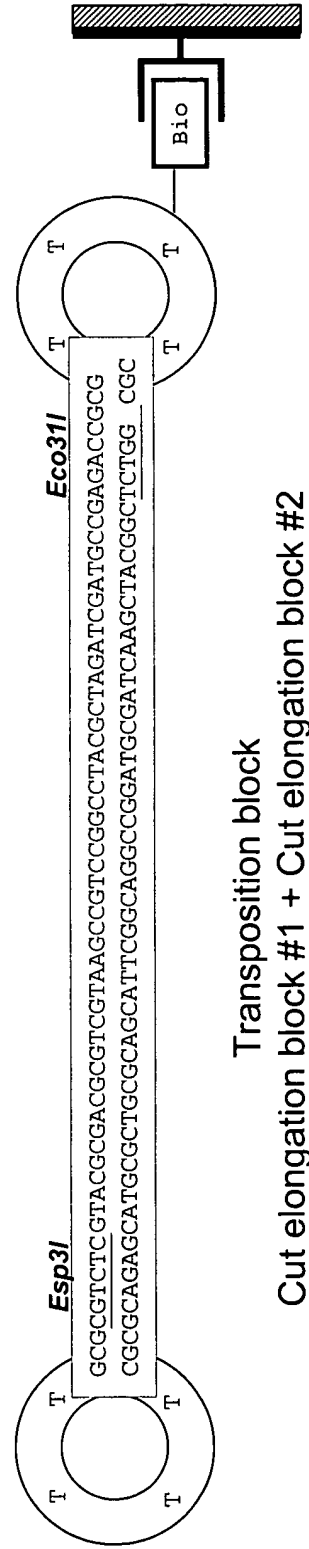
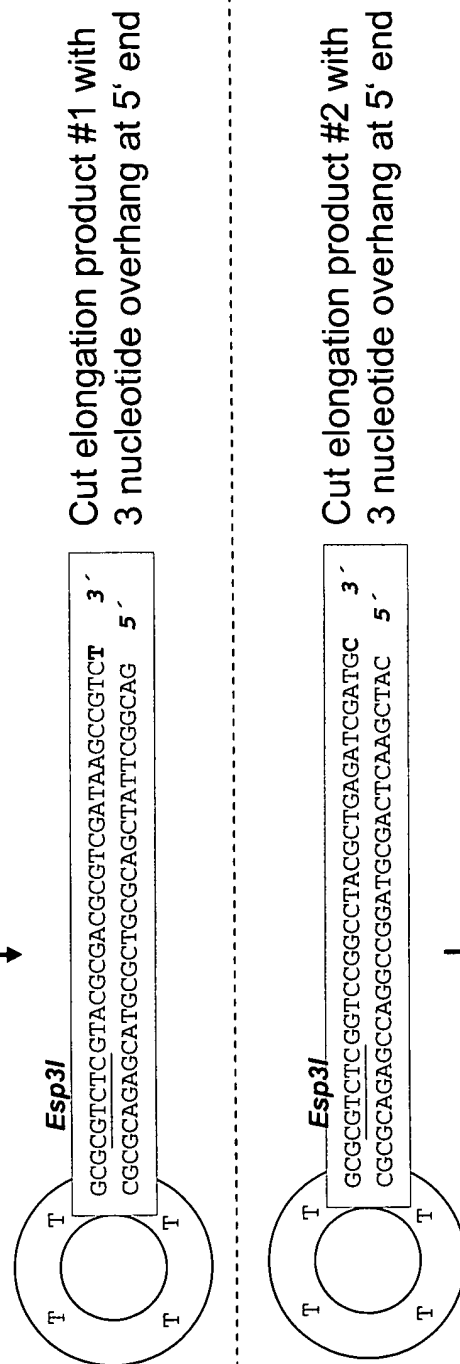
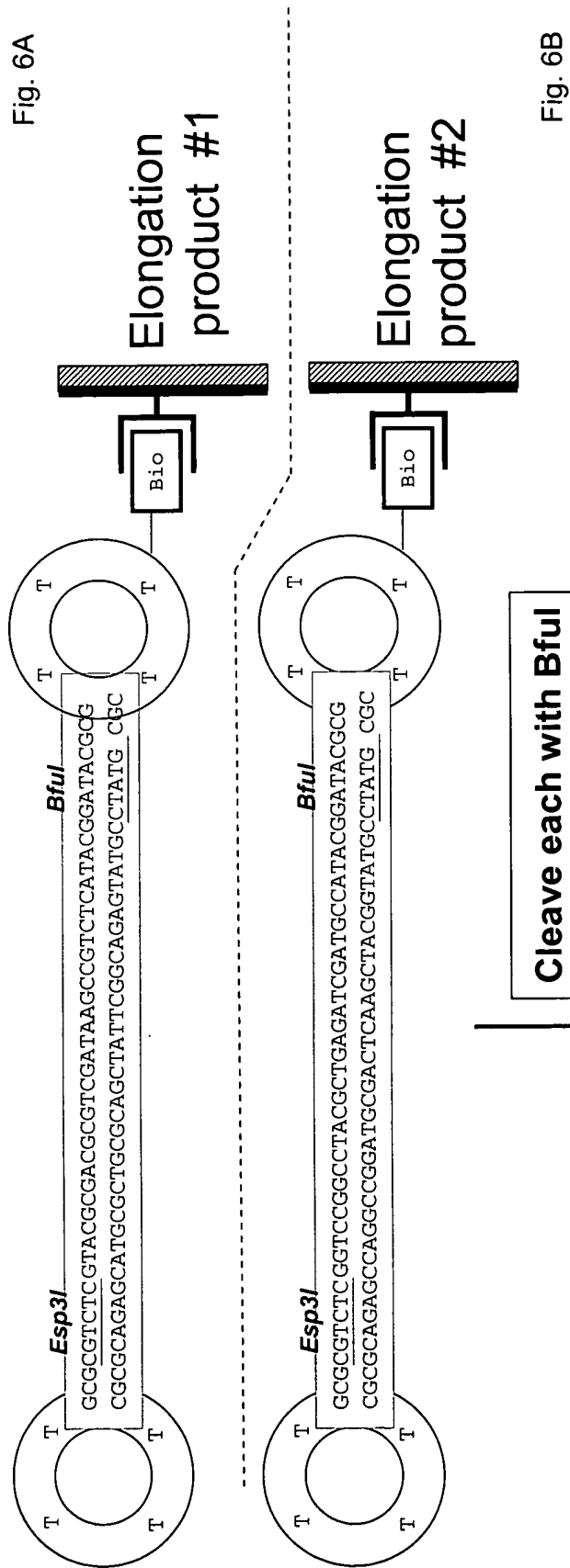


Fig. 5J



Transposition block
Cut elongation block #1 + Cut elongation block #2

Fig. 6 Addition of transition anchor (both RSPS and RLPs) and first transposition (1 nt overhang)



The diagram illustrates two transition anchors, Transition anchor #1 and Transition anchor #2, which are used for sequencing. Each anchor consists of a central DNA sequence flanked by two circular plasmid-like structures. The central DNA sequence is flanked by *Eco31I* and *Esp3I* restriction enzyme sites. The plasmid-like structures are labeled with 'T' and 'Bio' (BioLink).

Transition anchor #1

Central DNA sequence (5' to 3' on the top strand):

```

5' CGAGACCGCG
3' AGCTCTGGCGC

```

Flanking DNA sequence (5' to 3' on the top strand):

```

5' GCGGTCCTC GTACGCGACGCGTCGATAAGCCGTCT
3' CGGCAGAGCATGGCTGCGCAGCTATTTCGGCAG

```

Transition anchor #2

Central DNA sequence (5' to 3' on the top strand):

```

5' CGAGACCGCG
3' GGCTCTGGCGC

```

Flanking DNA sequence (5' to 3' on the top strand):

```

5' GCGGTCCTC GGTCCGGCCTACGCTGAGATCGATGC
3' CGGCAGAGCCAGGCCGGATGCGACTCAAGCTAC

```

Fig. 7 Semi-Inverted Transposition (SIT).

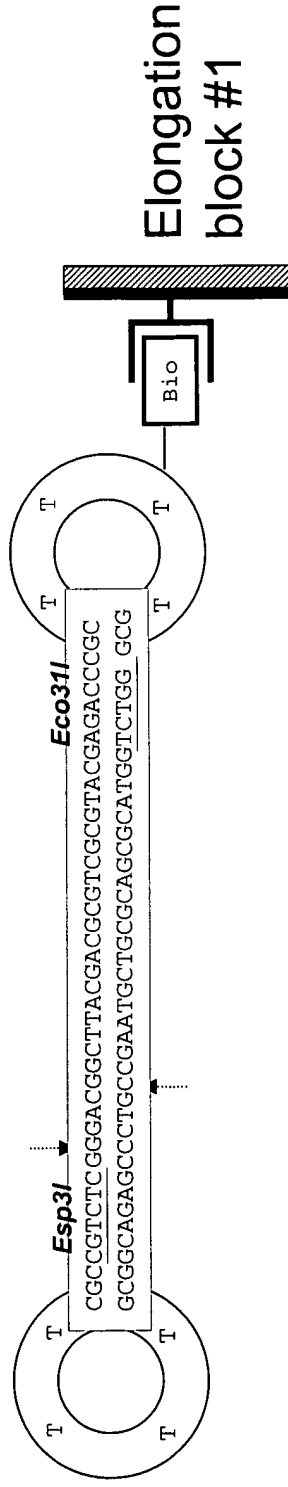


Fig. 7A

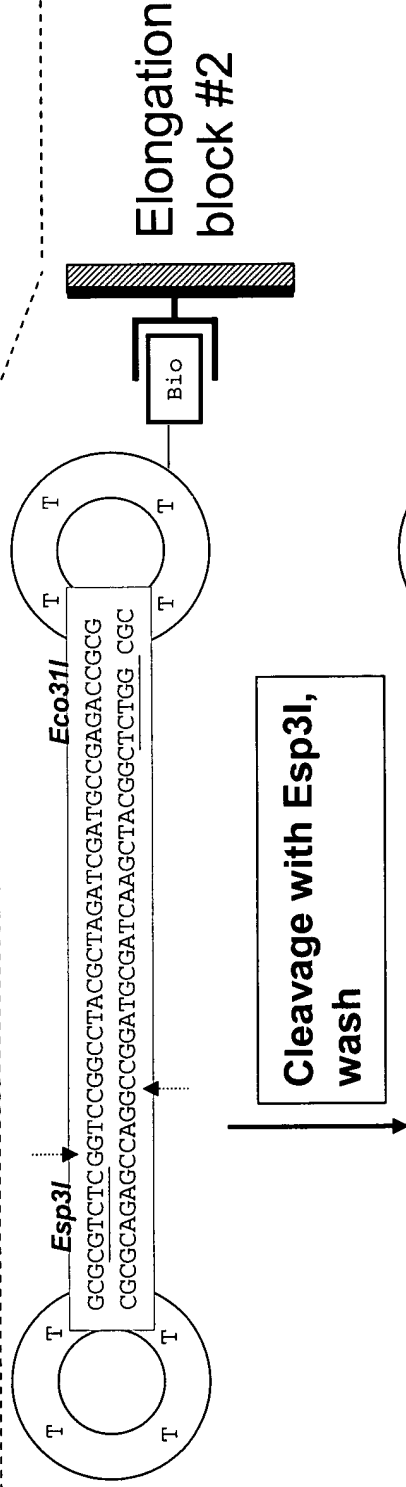
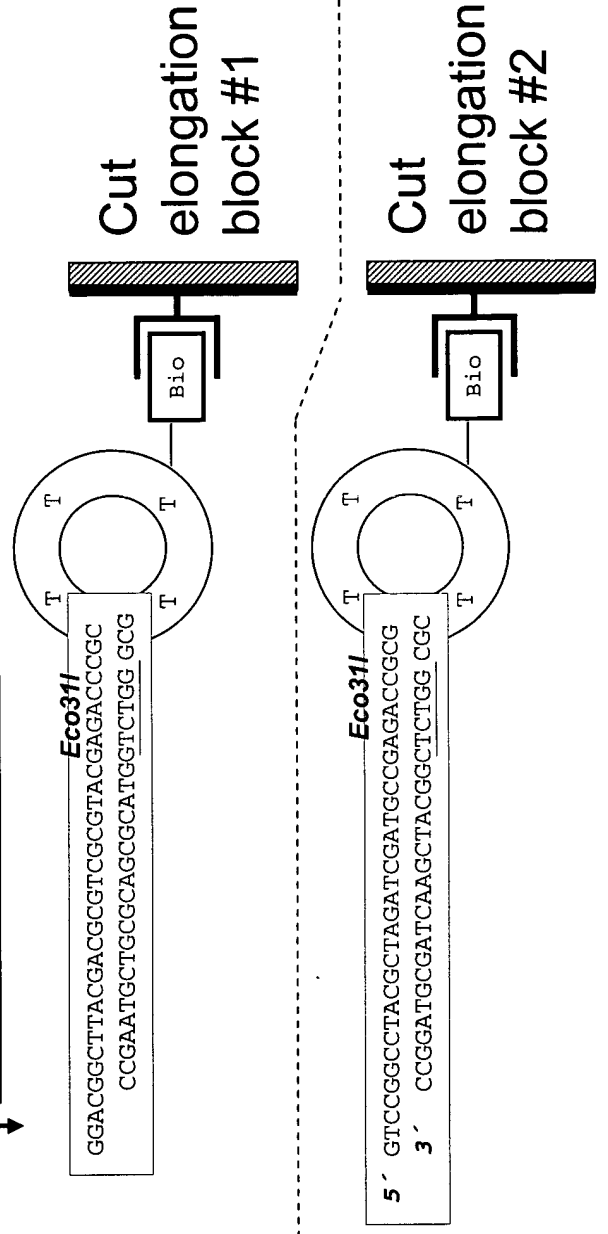


Fig. 7B



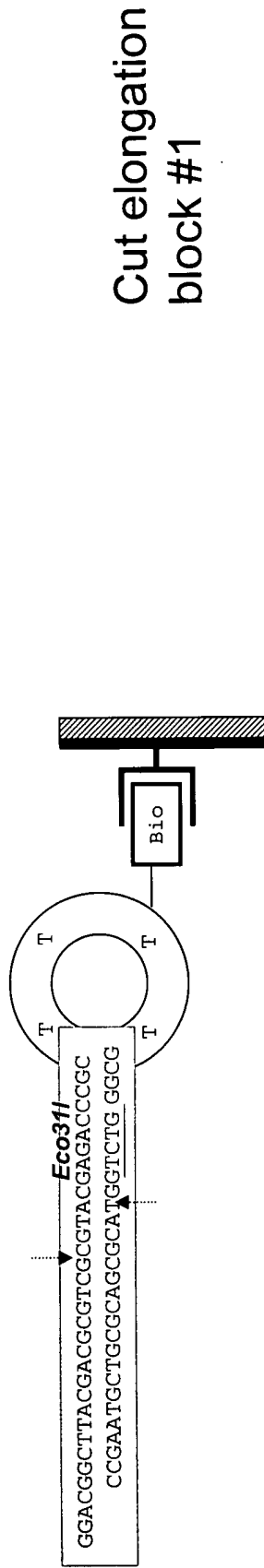


Fig. 7C

cleavage of elongation block #1 with *Eco31I*

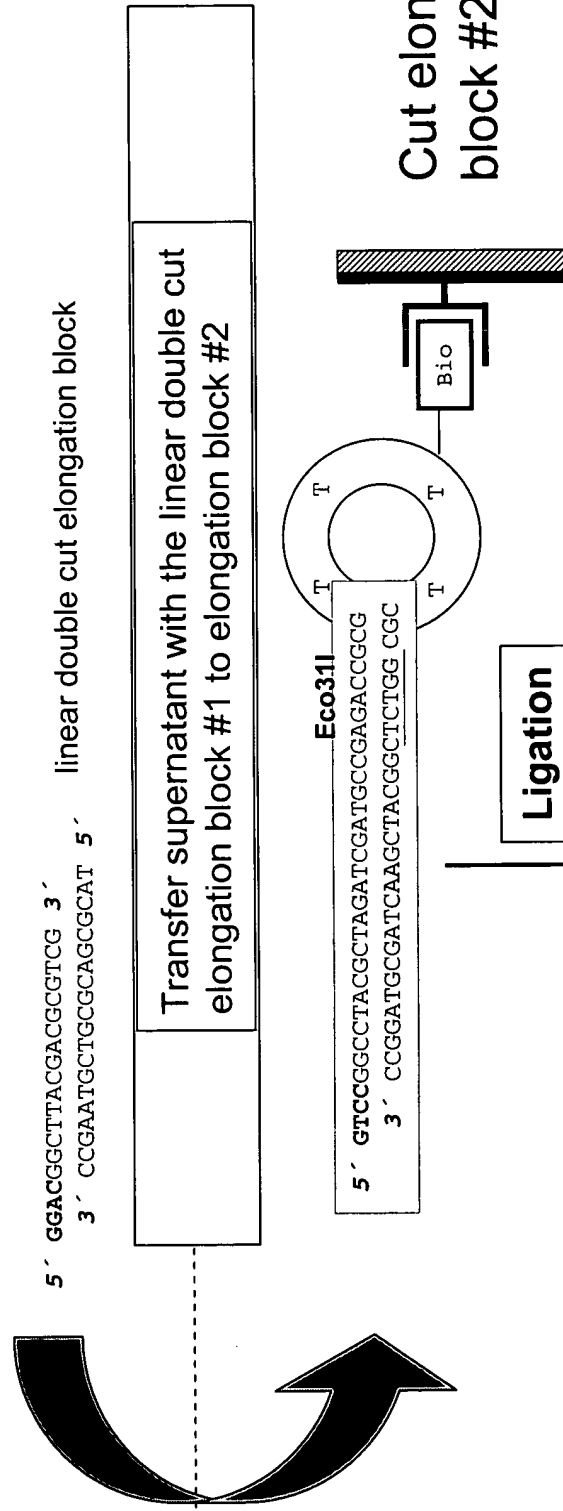
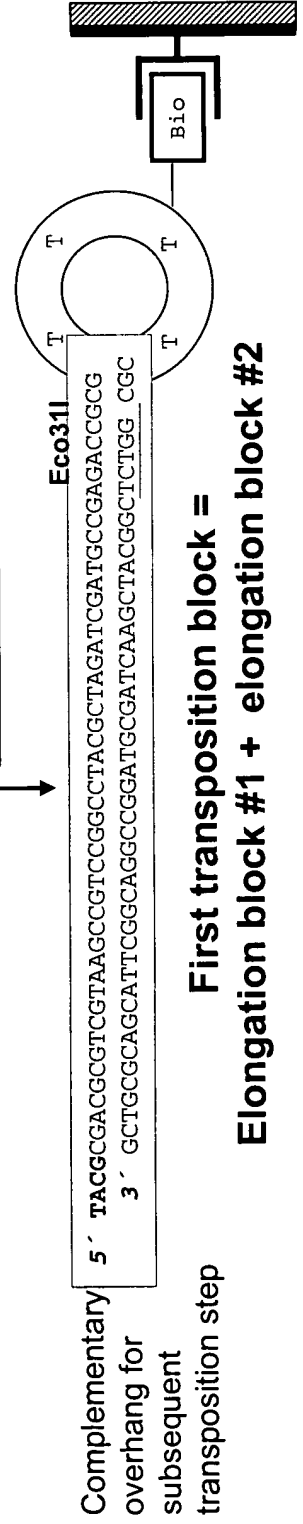


Fig. 7D

Ligation



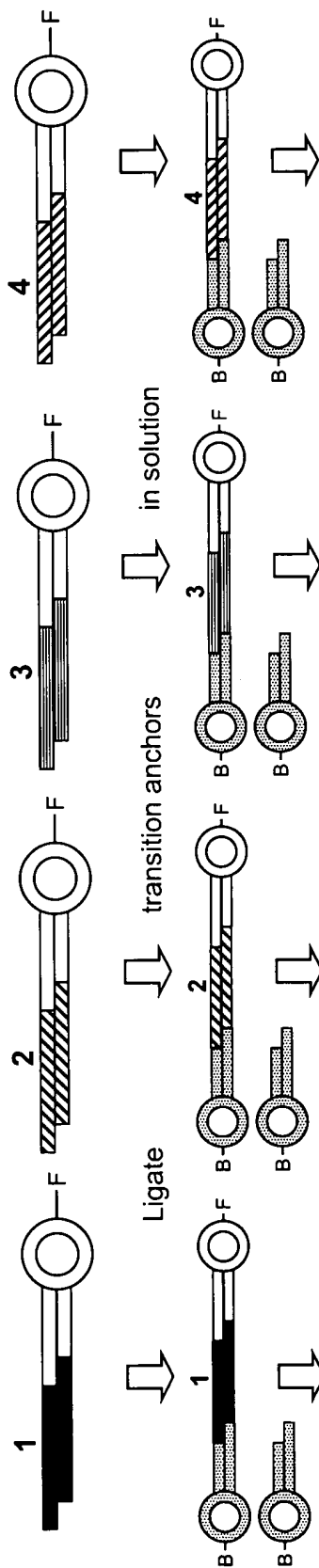
Complementary
overhang for
subsequent
transposition step

Double selection or pingpong procedure

Fig. 8A

Standard elongation reactions with FITC-labelled splinkers and Biotin-labelled anchors

Purified elongation products before addition of transition anchor (last further at least partially double-stranded oligonucleotide)



Immobilise on α -FITC, wash Immobilise on SA, wash Immobilise on α -FITC, wash Immobilise on SA, wash

Fig. 8B

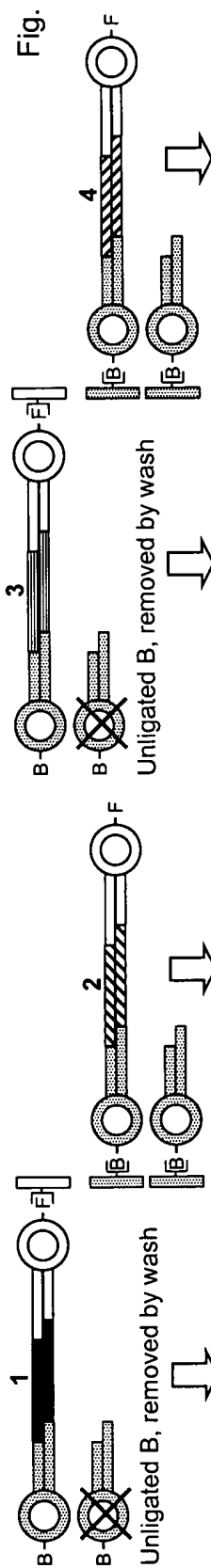


Fig. 8C

Cleave with Eco31I,
Transfer supernatant

Cleave with Esp3I,
Transfer supernatant

Cleave with Eco31I,
Transfer supernatant

Cleave with Esp3I,
Transfer supernatant

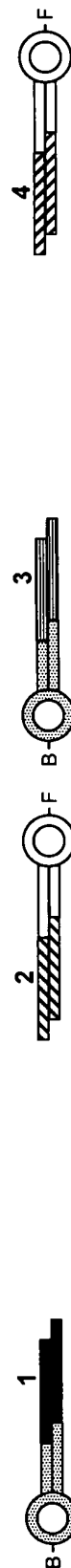


Fig. 8D



Double selection or pingpong procedure

Fig. 8E

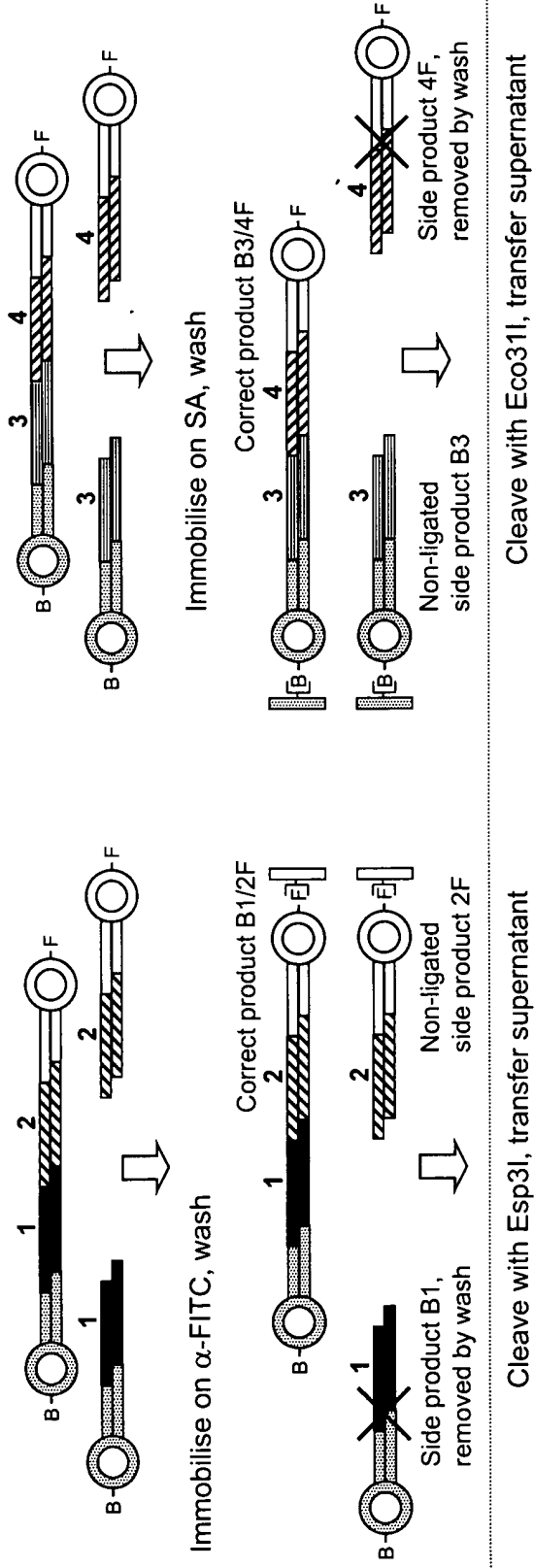
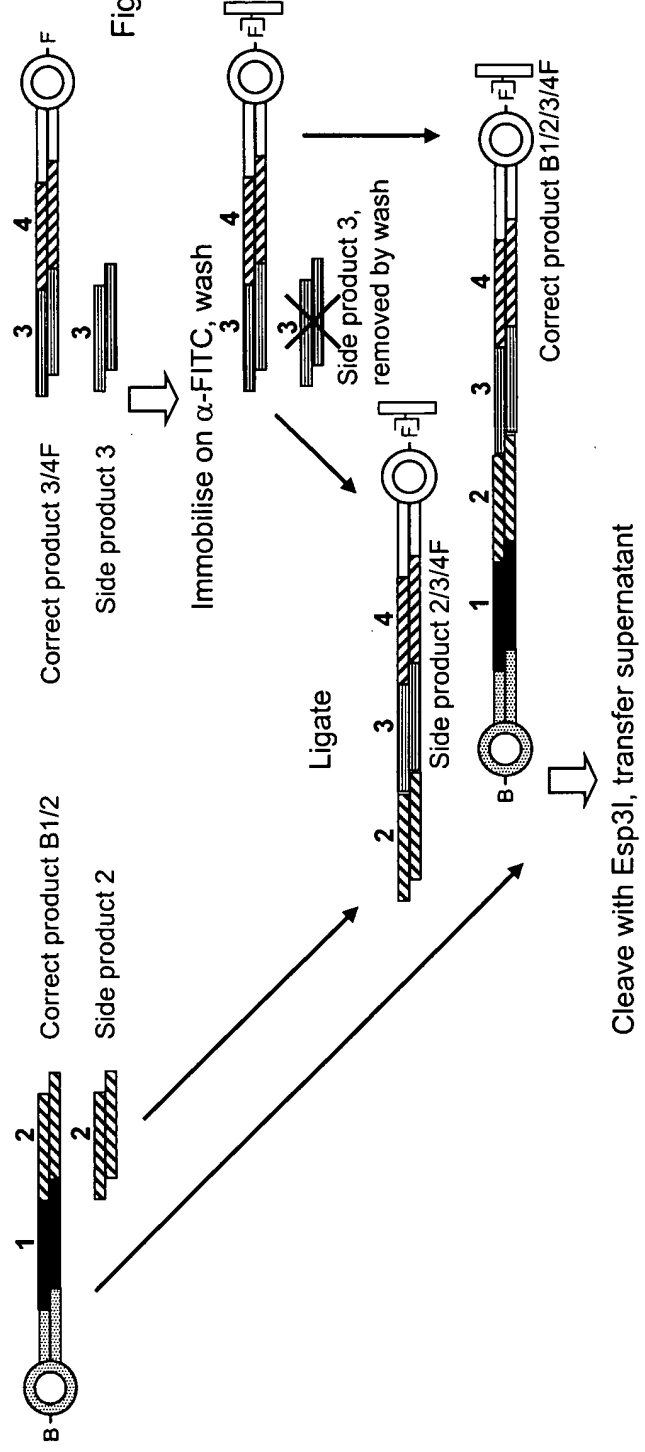
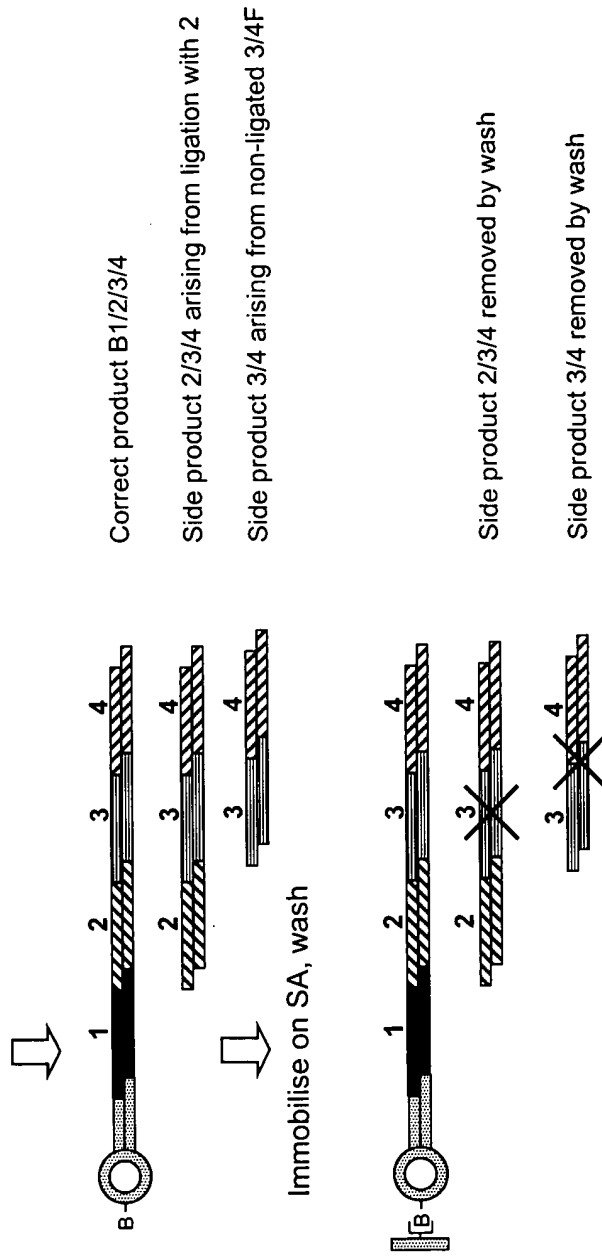


Fig. 8F



Double selection or pingpong procedure

Fig. 8G



10/531556

Semi-inverted transposition (SIT) with prior double selection

Fig. 9A

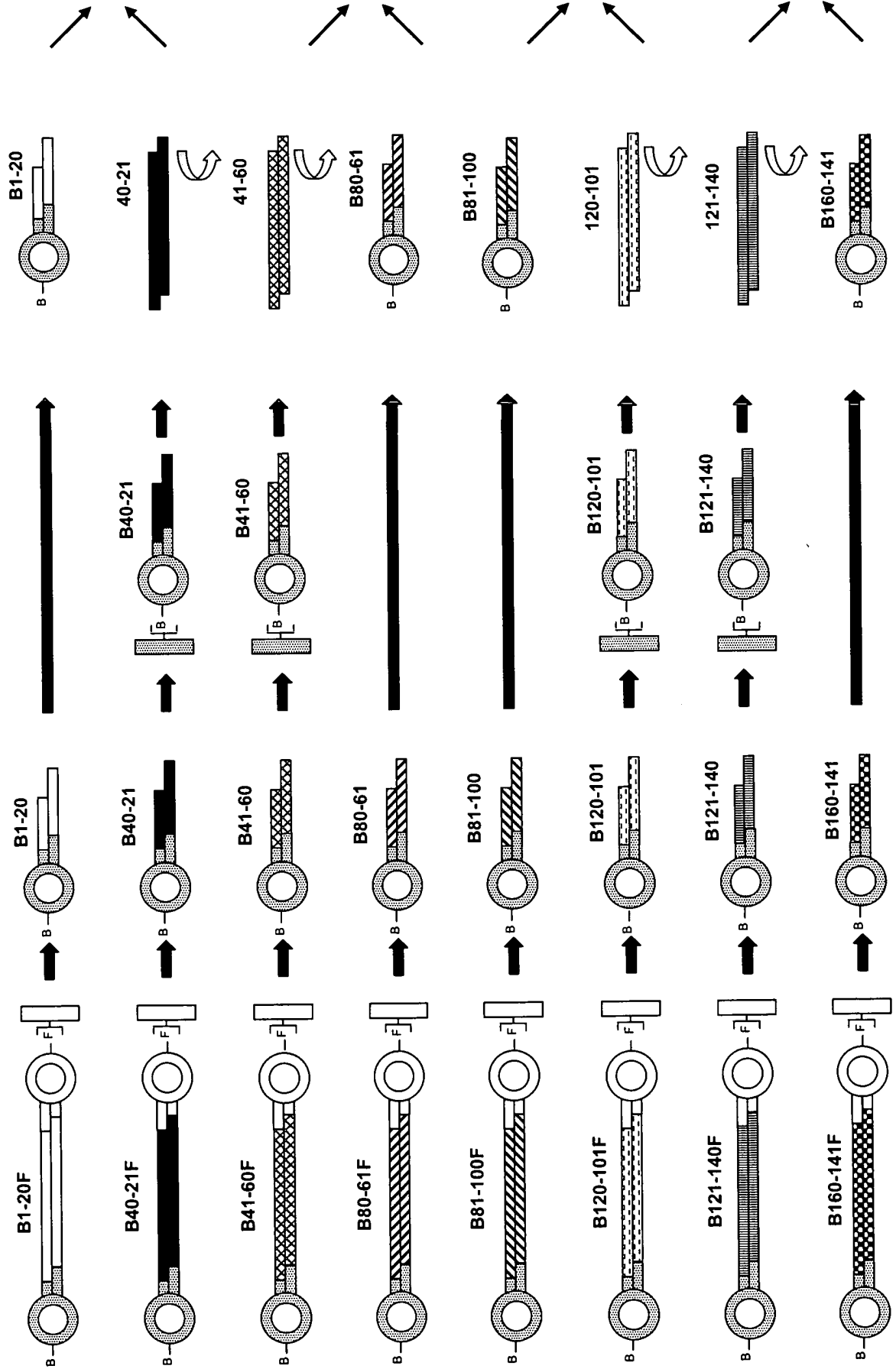
Immobilise elongation blocks via FITC

Cut all elongation blocks with RE1

Immobilise each other cut elongation block

Cut immobilised elongation blocks with RE2

Ligate pairs of single and double-cut elongation blocks

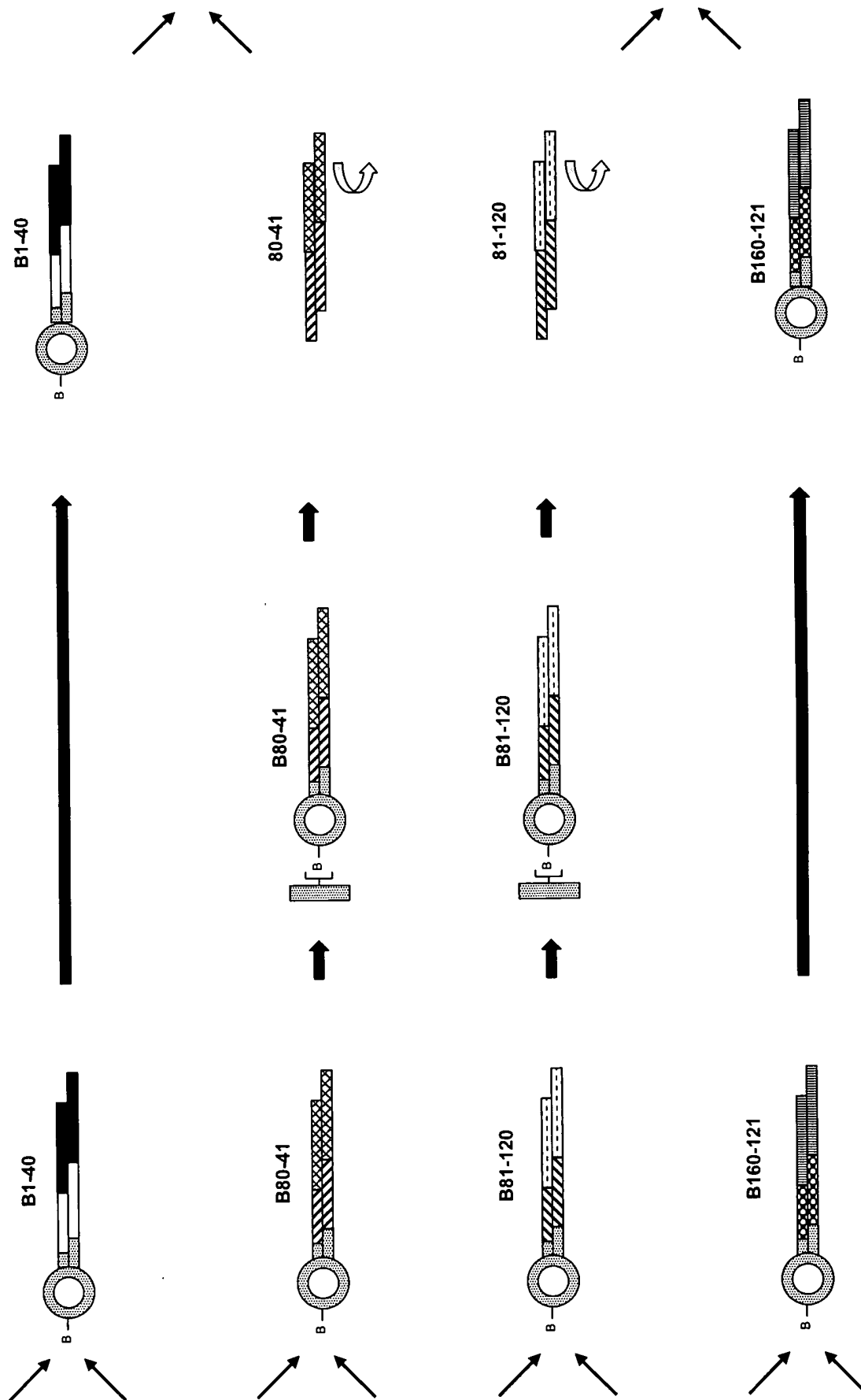


Semi-inverted transposition (SIT) with prior double selection

Fig. 9B

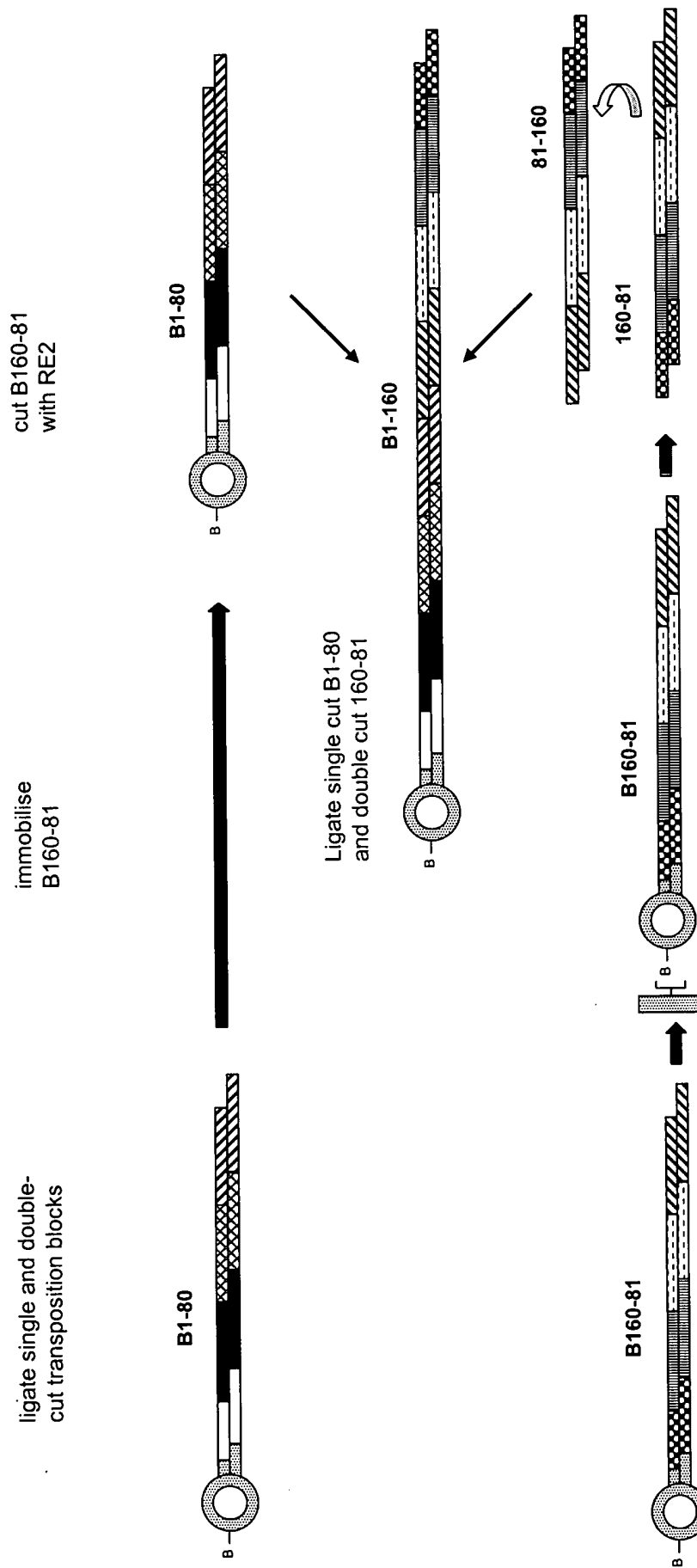
cut immobilised transposition blocks with RE2

immobilise alternate transposition blocks of each pair



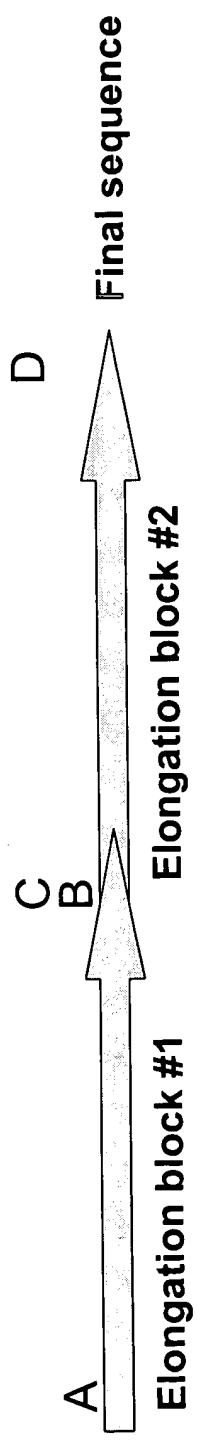
Semi-inverted transposition (SIT) with prior double selection

Fig. 9C

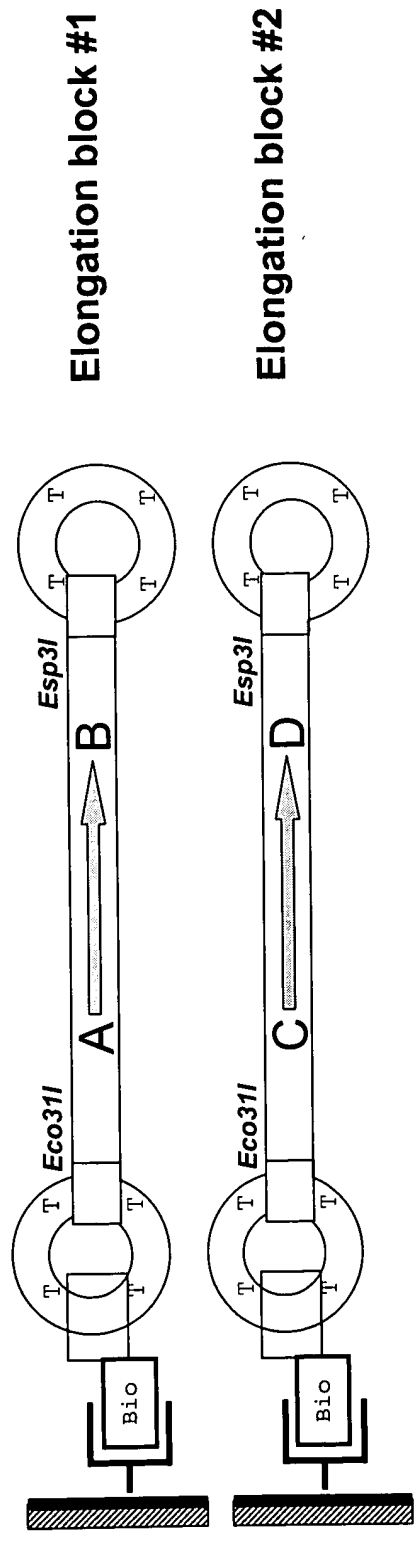


Design of elongation blocks for standard and semi-inverted transpositions

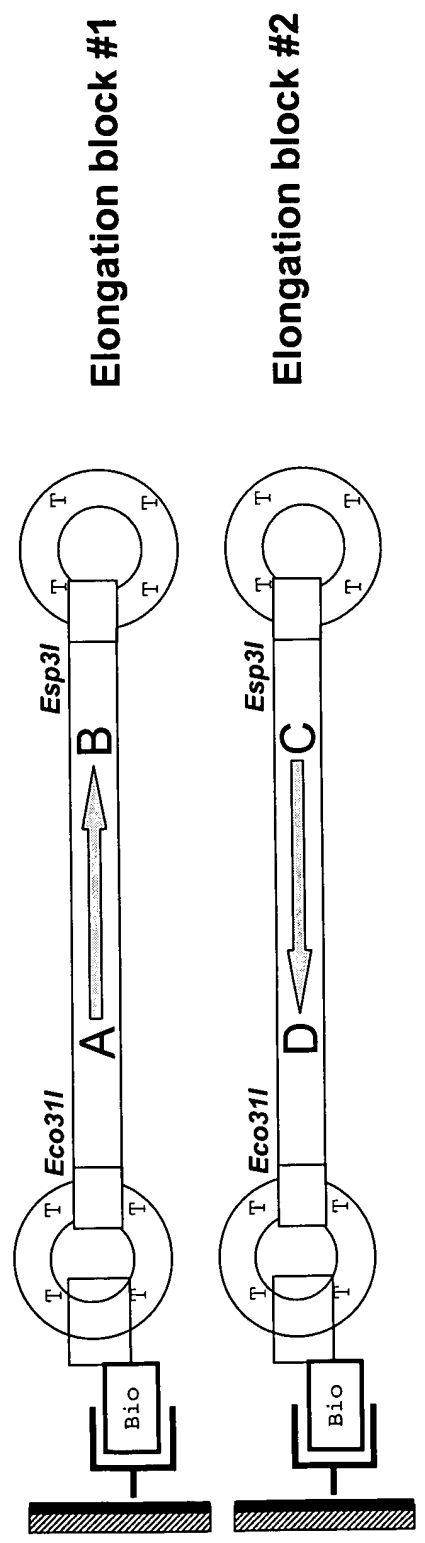
Fig. 10



Design of the Elongation blocks for Standard Transposition



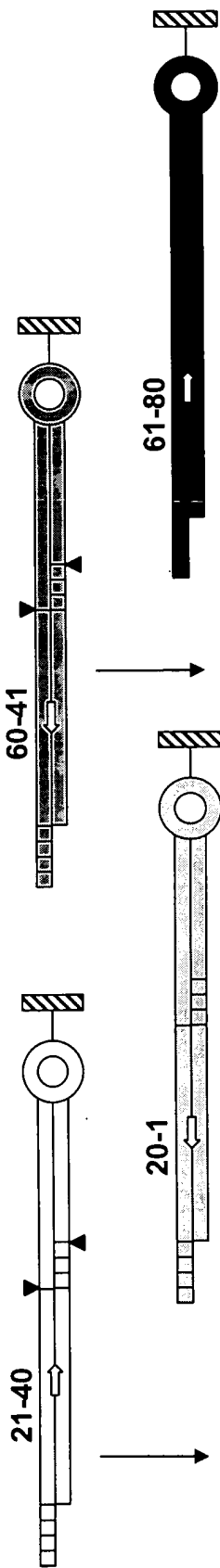
Design of the Elongation blocks for SIT



Semi-inverted transposition (SIT) with 3nt/4nt ligation

Fig. 11A

1. cleave all immobilised elongation blocks with RE specific for second at least partially double-stranded oligonucleotide



2. cleave every other cut immobilised elongation block with RE specific for further at least partially double-stranded oligonucleotide

Fig. 11B

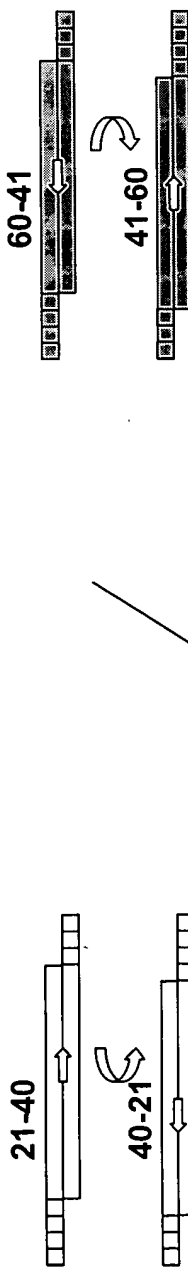
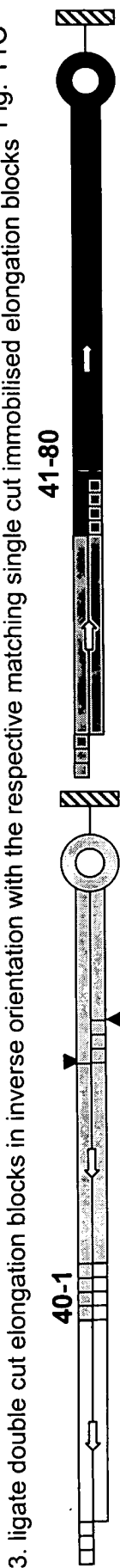


Fig. 11C



4. cleave every other immobilised transposition block with the same RE as before, ligate double cut transposition blocks in reverse orientation with their respective matching single cut immobilised transposition blocks

Fig. 11D

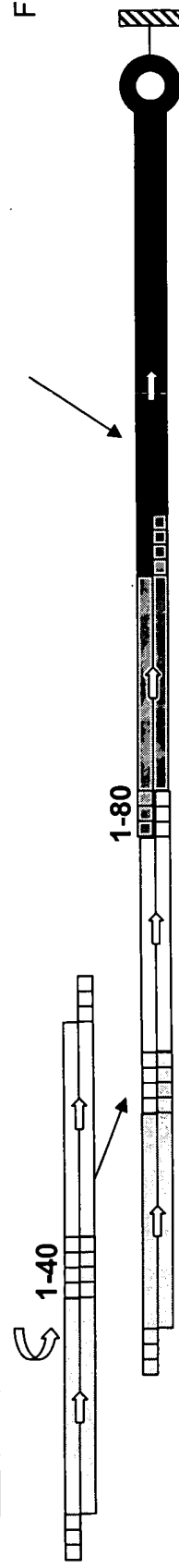
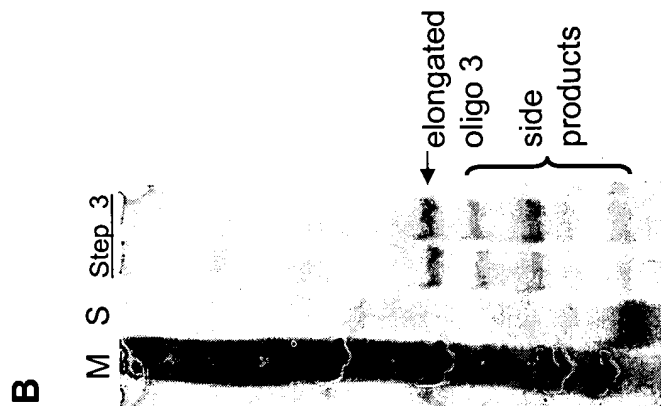
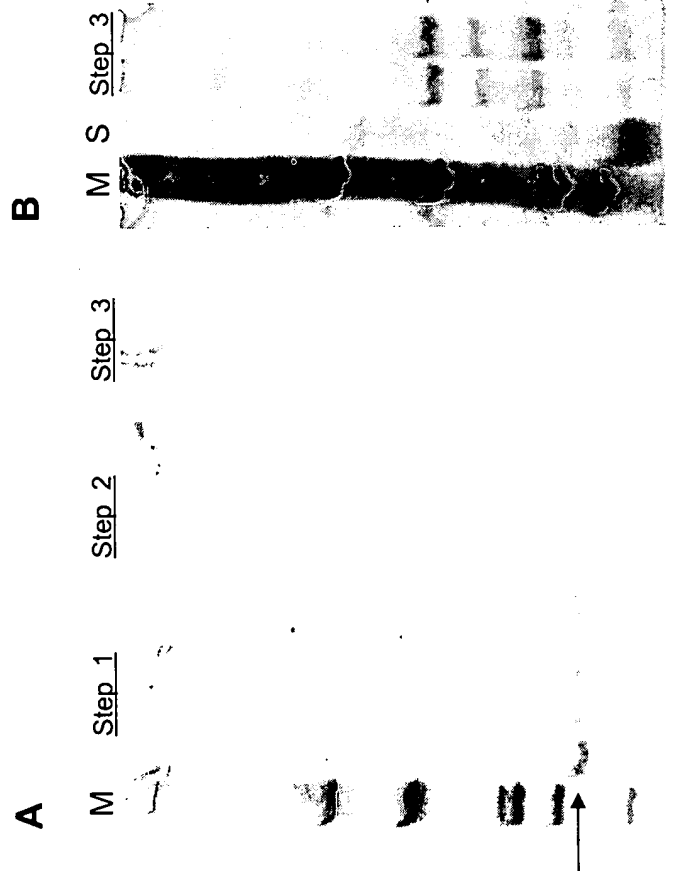
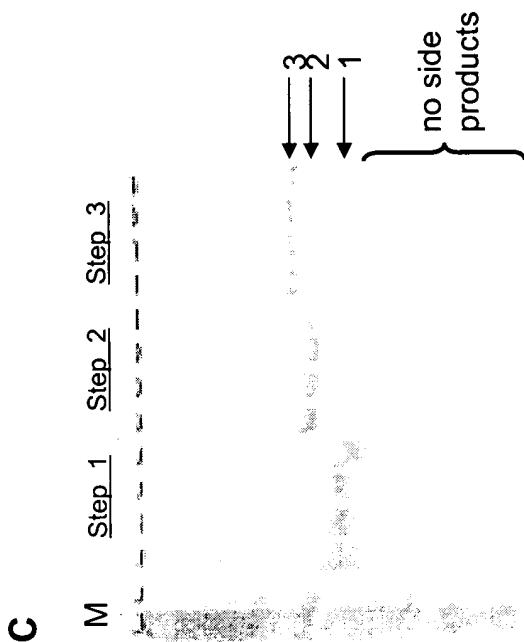


Fig. 12

SPS (solid phase synthesis)



RLPS (reverse liquid phase synthesis)



3: elongated oligo 3
2: elongated oligo 2
1: elongated oligo 1